SEQUENCE LISTING

<110> INCYTE PHARMACEUTICALS, INC.
HILLMAN, Jennifer L.; YUE, Henry
TANG, Y. Tom; AZIMZAI, Yalda

<120> CANCER ASSOCIATED PROTEINS

<130> PF-0661 USN

<140> US 09/889,617
<141> To Be Assigned

<150> US 00/01565
<151> 2000-01-21

<150> US 09/236,205
<151> 1999-01-22

<150> US 60/183,027
<151> 1999-01-22

<160> 9

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<212> PRT
<213> Homo sapiens

<220>
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Tyr Tyr Pro Asp Glu Ser Tyr Asn Glu Val Tyr Ala Glu Glu Val
             35                  40                  45
Pro Gln Ala Pro Ala Leu Asp Tyr Arg Val Pro Arg Trp Cys Tyr
             50                  55                  60
Thr Leu Asn Ile Gln Asp Gly Glu Ala Thr Cys Tyr Ser Pro Lys
             65                  70                  75
Gly Gly Asn Tyr His Ser Ser Leu Gly Thr Arg Cys Glu Leu Ser
             80                  85                  90
Cys Asp Arg Gly Phe Arg Leu Ile Gly Arg Arg Ser Val Gln Cys
             95                 100                 105
Leu Pro Ser Arg Arg Trp Ser Gly Thr Ala Tyr Cys Arg Gln Met
            110                 115                 120
Arg Cys His Ala Leu Pro Phe Ile Thr Ser Gly Thr Tyr Thr Cys
            125                 130                 135
Thr Asn Gly Val Leu Leu Asp Ser Arg Cys Asp Tyr Ser Cys Ser
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140 145 150
Ser Gly Tyr His Leu Glu Gly Asp Arg Ser Arg Ile Cys Met Glu
155 160 165
Asp Gly Arg Trp Ser Gly Gly Glu Pro Val Cys Val Asp Ile Asp
170 175 180
Pro Pro Lys Ile Arg Cys Pro His Ser Arg Glu Lys Met Ala Glu
185 190 195
Pro Glu Lys Leu Thr Ala Arg Val Tyr Trp Asp Pro Pro Leu Val
200 205 210
Lys Asp Ser Ala Asp Gly Thr Ile Thr Arg Val Thr Leu Arg Gly
215 220 225
Pro Glu Pro Gly Ser His Phe Pro Glu Gly Glu His Val Ile Arg
230 235 240
Tyr Thr Ala Tyr Asp Arg Ala Tyr Asn Arg Ala Ser Cys Lys Phe
245 250 255
Ile Val Lys Val Gln Val Arg Arg Cys Pro Thr Leu Lys Pro Pro
260 265 270
Gln His Gly Tyr Leu Thr Cys Thr Ser Ala Gly Asp Asn Tyr Gly
275 280 285
Ala Thr Cys Glu Tyr His Cys Asp Gly Gly Tyr Asp Arg Gln Gly
290 295 300
Thr Pro Ser Arg Val Cys Gln Ser Ser Arg Gln Trp Ser Gly Ser
305 310 315
Pro Pro Ile Cys Ala Pro Met Lys Ile Asn Val Asn Val Asn Ser
320 325 330
Ala Ala Gly Leu Leu Asp Gln Phe Tyr Glu Lys Gln Arg Leu Leu
335 340 345
Ile Ile Ser Ala Pro Asp Pro Ser Asn Arg Tyr Tyr Lys Met Gln
350 355 360
Ile Ser Met Leu Gln Gln Ser Thr Cys Gly Leu Asp Leu Arg His
365 370 375
Val Thr Ile Ile Glu Leu Val Gly Gln Pro Pro Gln Glu Val Gly
380 385 390
Arg Ile Arg Glu Gln Gln Leu Ser Ala Asn Ile Ile Glu Glu Leu
395 400 405
Arg Gln Phe Gln Arg Leu Thr Arg Ser Tyr Phe Asn Met Val Leu
410 415 420
Ile Asp Lys Gln Gly Ile Asp Arg Asp Arg Tyr Met Glu Pro Val
425 430 435
Thr Pro Glu Glu Ile Phe Thr Phe Ile Asp Asp Tyr Leu Leu Ser
440 445 450
Asn Gln Glu Leu Thr Gln Arg Arg Glu Gln Arg Asp Ile Cys Glu
455 460 465

<210> 2
<211> 400
<212> PRT
<213> Homo sapiens

<220>
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<223> Incyte ID No: 2616269CD1

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35 40 45
Leu Arg Arg Phe Thr Ser Cys Asp Leu Leu Trp Arg Arg Ile Ala
50 55 60
Arg Ala Ser Leu Asn Ser Gly Phe Thr Arg Leu Gly Thr Asp Leu
65 70 75
Met Thr Ser Val Pro Val Lys Glu Arg Val Lys Val Ser Gln Asn
80 85 90
Trp Arg Leu Gly Arg Cys Arg Glu Gly Ile Leu Leu Lys Trp Arg
95 100 105
Cys Ser Gln Met Pro Trp Met Gln Leu Glu Asp Asp Ser Leu Tyr
110 115 120
Ile Ser Gln Ala Asn Phe Ile Leu Ala Tyr Gln Phe Arg Pro Asp
125 130 135
Gly Ala Ser Leu Asn Arg Arg Pro Leu Gly Val Phe Ala Gly His
140 145 150
Asp Glu Asp Val Cys His Phe Val Leu Ala Asn Ser His Ile Val
155 160 165
Ser Ala Gly Gly Asp Gly Lys Ile Gly Ile His Lys Ile His Ser
170 175 180
Thr Phe Thr Val Lys Tyr Ser Ala His Glu Gln Glu Val Asn Cys
185 190 195
Val Asp Cys Lys Gly Gly Ile Ile Val Ser Gly Ser Arg Asp Arg
200 205 210
Thr Ala Lys Val Trp Pro Leu Ala Ser Gly Arg Leu Gly Gln Cys
215 220 225
Leu His Thr Ile Gln Thr Glu Asp Arg Val Trp Ser Ile Ala Ile
230 235 240
Ser Pro Leu Leu Ser Ser Phe Val Thr Gly Thr Ala Cys Cys Gly
245 250 255
His Phe Ser Pro Leu Arg Ile Trp Asp Leu Asn Ser Gly Gln Leu
260 265 270
Met Thr His Leu Gly Ser Asp Phe Pro Pro Gly Ala Gly Val Leu
275 280 285
Asp Val Met Tyr Glu Ser Pro Phe Thr Leu Leu Ser Cys Gly Tyr
290 295 300
Asp Thr Tyr Val Arg Tyr Trp Asp Leu Arg Thr Ser Val Arg Lys
305 310 315
Cys Val Met Glu Trp Glu Glu Pro His Asp Ser Thr Leu Tyr Cys
320 325 330
Leu Gln Thr Asp Gly Asn His Leu Leu Ala Thr Gly Ser Ser Tyr
335 340 345
Tyr Gly Val Val Arg Leu Trp Asp Arg Arg Gln Arg Ala Cys Leu
350 355 360
His Ala Phe Pro Leu Thr Ser Thr Pro Leu Ser Ser Pro Val Tyr
365 370 375
Cys Leu Arg Leu Thr Thr Lys His Leu Tyr Ala Ala Leu Ser Tyr
380 385 390
Asn Leu His Val Leu Asp Phe Gln Asn Pro
395 400

<210> 3

<211> 146
<212> PRT
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 3117642CD1

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35 40 45
Arg Glu Asp Glu Gly Ala Ser Ala Ala Gln Gly Gln Glu Pro Glu
50 55 60
Ala Asp Ser Gln Glu Leu Val Gln Pro Lys Thr Gly Cys Glu Leu
65 70 75
Gly Asp Gly Pro Asp Thr Lys Arg Val Cys Leu Arg Asn Glu Glu
80 85 90
Gln Met Lys Leu Pro Ala Glu Gly Pro Glu Pro Glu Ala Asp Ser
95 100 105
Gln Glu Gln Val His Pro Lys Thr Gly Cys Glu Arg Gly Asp Gly
110 115 120
Pro Asp Val Gln Glu Leu Gly Leu Pro Asn Pro Glu Glu Val Lys
125 130 135
Thr Pro Glu Glu Asp Glu Gly Gln Ser Gln Pro
140 145

<210> 4
<211> 2152
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 1518859CB1

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ttggcagatg tggacgtcgt aacatctggg cagtgttaac agaatcccgg aggccgggac 180
agaccaggag ccactcgttc taggaatgtt aaagtagaag gttttttcca attgatgaga 240
ggagcagaga ggaaggagaa agaggaggag agagaaaaag ggcacaaaat accataaaac 300
agatcccata tttctgcttc ccctcacttt tagaagttaa ttgatggctg acttctgaaa 360
gtcactttcc tttgccctgg tacttcaggc catatacatc ttttcttgtc tccataatcc 420
tccctttcaa ggatggccag tcagctaact caaagaggag ctctctttct gctgttcttc 480
ctaactccgg cagtgacacc aacatggtat gcaggttctg gctactatcc ggatgaaagc 540
tacaatgaag tatatgcaga ggaggtccca caggctcctg ccctggacta ccgagtcccc 600
cgatggtgtt atacattaaa tatccaggat ggagaagcca catgctactc accgaaggga 660
ggaaattatc acagcagcct gggcacgcgt tgtgagctct cctgtgaccg gggctttcga 720
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tgcaggcaga tgagatgcca cgcactacca ttcatcacta gtggcactta cacctgcaca 840
aatggagtgc ttcttgactc tcgctgtgac tacagctgtt ccagtggcta ccacctggaa 900

ggtgatcgca gccgaatctg catggaagat gggagatgga gtggaggcga gcctgtatgt 960
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gagaaattga ctgctcgagt atactgggac ccaccgttgg tgaaagattc tgctgatggt 1080
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cccgaggaaa tcttcacatt cattgatgac tacctactga gcaatcagga gttgacccag 1800
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agttctaggg acaggactct gaggtgggtg agtttgacaa atcctgcagt gtttccaggc 1980
atcctttag gactgtgtaa tagtttccct agaagctagg tagggactga ggacaggcct 2040
tgggcagtgg gttgggggta gaagttcttc ctttcctaac ccgggcccct gcccagctct 2100
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<210> 5
<211> 1888
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 2616269CB1

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tcatctgctc ctacctggac atgcgggccc tcggccgcct ggcccaggtg tgccgctggc 180
tgcggcgctt caccagctgc gatctgctct ggcgccggat agcccgggcc tcgctcaact 240
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aggtgtctca gaactggaga ctggggcgct gccgagaggg gattctgctg aagtggagat 360
gcagtcagat gccctggatg cagctagagg atgattctct gtacatatcc caggctaatt 420
tcatcctggc ctaccagttc cgtccagatg gtgccagctt gaatcgtcgg cctctgggag 480
tctttgctgg gcatgatgag gacgtttgcc actttgtgct ggccaactcg catattgtta 540
gtgcaggagg ggatgggaag attggcattc ataagattca cagcaccttc actgtcaagt 600
actcggctca tgaacaggag gtgaactgtg tggattgcaa agggggcatc attgtgagtg 660
gctccaggga caggacggcc aaggtgtggc ctttggcctc aggccggctg gggcagtgct 720
tacacaccat ccagactgaa gaccgagtct ggtccattgc tatcagccca ttactcagct 780
cttttgtgac agggacggct tgttgcgggc acttctcacc cctgagaatc tgggacctca 840
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atgtcatgta tgagtccccc ttcacactgc tgtcctgtgg ctatgacacc tatgttcgct 960
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ctgccctgtc ttacaacctc cacgtcctgg attttcaaaa cccatgaccg tcagggccac 1260
ccctgcctct gggccaggga aaccagctac tcagggactt ctcttgcctg gagggtgcag 1320
tgatagctcc tcctcactgc cccactgtgc tcctgggcct gtgaccccag tgctcaggca 1380

ccttgcacta gaggcttctg actcctggga ctttggagct taccagagat gcagtccctc 1440
ccaggaacct gttggagagg caggacctgc tgctttagag tgcggctgaa cccgggcctt 1500
gcatccctgt ttggccagag caaggatctg gcctggagag gcccatccta tacccctttat 1560
tamagccatg acagcctaca gagtgaggtg aggtgctccc accttcccag atggttcctt 1620
tctgcccctt cctggaagga aaggtgaggc tgccaatagc ctcctggcac cagccagacc 1680
tcacccttga ccaacctctc ggggctgggg gttcattcct ggggcactgt ggcctggttt 1740
tgctttgaaa ccaagaaaga gcaaagggaa cccagcagtt ctgagtgagt tctgagccag 1800
ccctacctca ggctggctgt tgagacatgc tacaattttc atttttgtaa aaataaagct 1860
tgattgttca cagaaaaaaa aaaaaaaa 1888

<210> 6
<211> 650
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 3117642CB1

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ctaagaagat taatctatcg gcgtagacca atgatctatg tagaatcttc tgaggagtcc 180
agtgatgagc aacctgacga agtggaatca ccaactcaaa gtcaggattc tacacctgct 240
gaagagagag aggatgaggg agcatctgca gctcaagggc aggagcctga agctgatagc 300
caggaactgg ttcagccaaa gactgggtgt gagcttggag atggtcctga taccaagagg 360
gtgtgcctgc gaaatgaaga gcagatgaaa ctgcccgcag aagggccaga gcctgaagcg 420
gatagccagg aacaggttca cccgaagact gggtgtgagc gcggagatgg tcctgatgtc 480
caggagttgg gcctgccaaa tccagaggag gtgaaaacac ctgaggaaga tgaagggcaa 540
tcacagcctt aaaagaagac acgctgaaat ggttcaggct gctcctgtgt tggaaatttg 600
accattaaaa ttctcccaat aaagctttac agccttctgc aaaaaaaaaa 650

<210> 7
<211> 464
<212> PRT
<213> Rattus norvegicus

<220>
<221> misc_feature
<223> GenBank ID No: g1345423

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Phe Pro Gly Ser Gly Asp Ser Pro Leu Glu Asp Asp Gly Val Trp
35 40 45
Ser Ser His Ser Leu Tyr Lys Asp Thr Pro Trp Cys Ser Pro Ile
50 55 60
Lys Val Lys Tyr Gly Asp Val Tyr Cys Arg Ala Pro Pro Gly Gly
65 70 75
Tyr Tyr Lys Thr Ala Leu Gly Thr Arg Cys Asp Ile Arg Cys Arg
80 85 90

Lys Gly Tyr Glu Leu His Gly Ser Ser Gln Leu Val Cys Gln Ser
95 100 105
Asn Arg Arg Trp Ser Asp Lys Val Ile Cys Lys Gln Lys Arg Cys
110 115 120
Pro Thr Leu Thr Met Pro Ala Asn Gly Gly Phe Lys Cys Val Asp
125 130 135
Gly Ala Tyr Phe Asn Ser Arg Cys Glu Tyr Tyr Cys Ser Pro Gly
140 145 150
Tyr Thr Leu Lys Gly Glu Arg Thr Val Thr Cys Met Asp Asn Lys
155 160 165
Ala Trp Ser Gly Arg Pro Ala Ser Cys Val Asp Met Glu Pro Pro
170 175 180
Arg Ile Lys Cys Pro Ser Val Lys Glu Arg Ile Ala Glu Pro Asn
185 190 195
Lys Leu Thr Val Arg Val Ser Trp Glu Thr Pro Glu Gly Arg Asp
200 205 210
Thr Ala Asp Gly Ile Leu Thr Asp Val Ile Leu Arg Gly Leu Pro
215 220 225
Pro Gly Ser Asn Phe Pro Glu Gly Asp His Lys Ile Glu Tyr Thr
230 235 240
Val Tyr Asp Arg Ala Glu Asn Lys Gly Thr Cys Lys Phe Arg Val
245 250 255
Lys Val Arg Val Arg Arg Cys Gly Lys Leu Asn Ala Pro Glu Asn
260 265 270
Gly Tyr Met Lys Cys Ser Ser Asp Gly Asp Asn Tyr Gly Ala Thr
275 280 285
Cys Glu Phe Ser Cys Ile Gly Gly Tyr Glu Leu Gln Gly Ser Pro
290 295 300
Ala Arg Val Cys Gln Ser Asn Leu Ala Trp Ser Gly Thr Glu Pro
305 310 315
Ser Cys Ala Ala Met Asn Val Asn Val Gly Val Arg Thr Ala Ala
320 325 330
Ala Leu Leu Asp Gln Phe Tyr Glu Lys Arg Arg Leu Leu Ile Val
335 340 345
Ser Thr Pro Thr Ala Arg Asn Leu Leu Tyr Arg Leu Gln Leu Gly
350 355 360
Met Leu Gln Gln Ala Gln Cys Gly Leu Asp Leu Arg His Ile Thr
365 370 375
Val Val Glu Leu Val Gly Val Phe Pro Thr Leu Ile Gly Arg Ile
380 385 390
Arg Ala Lys Ile Met Pro Pro Ala Leu Ala Leu Gln Leu Arg Leu
395 400 405
Leu Leu Arg Ile Pro Leu Tyr Ser Phe Ser Met Val Leu Val Asp
410 415 420
Lys His Gly Met Asp Lys Glu Arg Tyr Val Ser Leu Val Thr Pro
425 430 435
Met Ala Leu Phe Asn Leu Ile Asp Thr Phe Pro Leu Arg Lys Glu
440 445 450
Glu Met Ile Leu Gln Ala Glu Met Gly Gln Ser Cys Asn Thr
455 460

<210> 8
<211> 278
<212> PRT
<213> Homo sapiens

<220>
<221> misc_feature
<223> GenBank ID No: g487348

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35 40 45
Ala Arg Thr Arg Pro Arg Glu Glu Ala Glu Gly Gly Gly Ser Val
50 55 60
Glu Glu Gly Ala Arg Gly Ile Ile Lys Gly Asp Glu Gly Ser Val
65 70 75
Gly Ala Gly Lys Glu Ala Gln Gly Arg Lys Tyr Gly Lys Glu Glu
80 85 90
Trp Arg Val Arg Ala Arg Arg Arg Glu Gly Ala Arg Pro Gly Arg
95 100 105
Val Gln Gly Gln Gly Gly Gln Val Trp Ala Tyr Ile Pro Gly Thr
110 115 120
Gly Ala Ala Met Ala Ala Ala Ala Arg Glu Glu Glu Glu Glu Ala
125 130 135
Ala Arg Glu Ser Ala Ala Cys Pro Ala Ala Gly Pro Ala Leu Trp
140 145 150
Arg Leu Pro Glu Val Leu Leu Leu His Met Cys Ser Tyr Leu Asp
155 160 165
Met Arg Ala Leu Gly Arg Leu Ala Gln Val Tyr Arg Trp Leu Trp
170 175 180
His Phe Thr Asn Cys Asp Leu Leu Arg Arg Gln Ile Ala Trp Ala
185 190 195
Ser Leu Asn Ser Gly Phe Thr Arg Leu Gly Thr Asn Leu Met Thr
200 205 210
Ser Val Pro Val Lys Val Ser Gln Asn Trp Ile Val Gly Cys Cys
215 220 225
Arg Glu Gly Ile Leu Leu Lys Trp Arg Cys Ser Gln Met Pro Trp
230 235 240
Met Gln Leu Glu Asp Asp Ala Leu Tyr Ile Ser Gln Ala Asn Phe
245 250 255
Ile Leu Ala Tyr Gln Phe Arg Pro Asp Gly Ala Ser Leu Asn Arg
260 265 270
Gln Pro Leu Gly Val Cys Trp Ala
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<210> 9
<211> 116
<212> PRT
<213> Homo sapiens

<220>
<221> misc_feature
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            20                  25                  30
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        35                  40                  45
Ala Thr Gln Arg Gln Asp Pro Ala Ala Ala Gln Glu Gly Glu Asp
    50                  55                  60
Glu Gly Ala Ser Ala Gly Gln Gly Pro Lys Pro Glu Ala Asp Ser
65                  70                  75
Gln Glu Gln Gly His Pro Gln Thr Gly Cys Glu Cys Glu Asp Gly
                80                  85                  90
Pro Asp Gly Gln Glu Met Asp Pro Pro Asn Pro Glu Glu Val Lys
            95                  100                 105
Thr Pro Glu Glu Gly Glu Lys Gln Ser Gln Cys
        110                 115
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